

Supporting Information Figure S3

Revised timeline and distribution of the earliest diverged human maternal lineages in southern Africa

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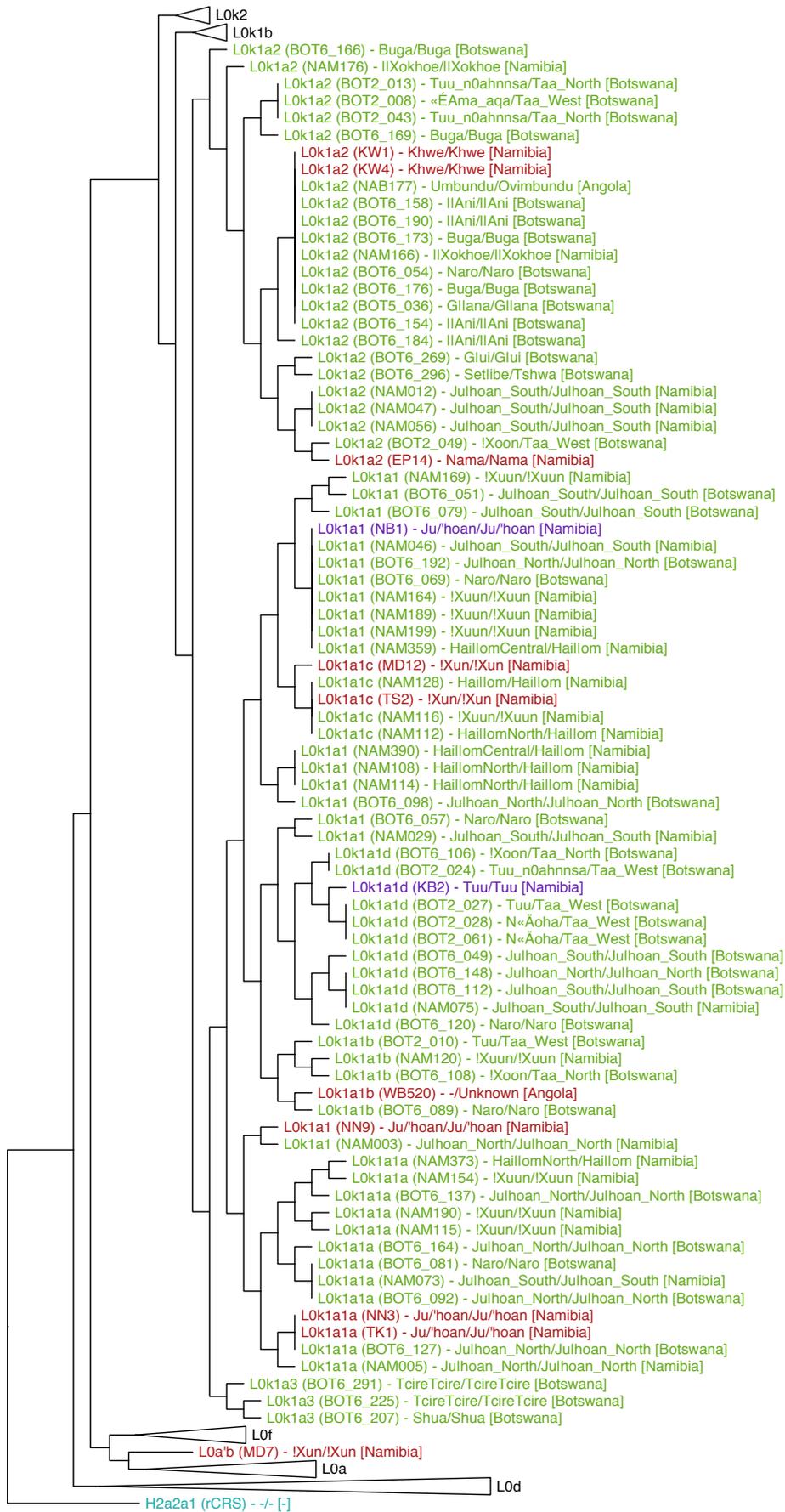


Fig. S3. Phylogenetic tree for L0k. Phylogeny was inferred using a total of 209 mtDNA, including all 139 mtDNA in the focus set and 70 additional L0k mtDNA from Barbieri *et al.* 2013. Haplogroups other than L0k are “collapsed”. Tips of L0k genomes are labeled with the format: Haplogroup (Isolate Name) - Language/Isolate Source [Country], where Isolate Name, Isolate Source, and Country are information included in the corresponding GenBank entries, and Language is the reported spoken language. Tip colours reflect data source: Red = current study, Green = Barbieri *et al.* 2013, Purple = Schuster *et al.* 2010, and Aqua = NCBI.